STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 0/821, 00/4Source: 0.3/0.3/2.006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



DATE: 03/03/2006

TIME: 15:07:14

IFW16

```
Input Set : A:\NS104D1C1 seqlisting rev1.txt
                    Output Set: N:\CRF4\03032006\J821001A.raw
      4 <110> APPLICANT: Palese, Peter
        Garcia-Sastre, Adolfo
      7 <120> TITLE OF INVENTION: RECOMBINANT NEGATIVE STRAND RNA VIRUS
             EXPRESSION SYSTEMS AND VACCINES
     11 <130> FILE REFERENCE: 26-003700US
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/821,001A
C--> 14 <141> CURRENT FILING DATE: 2004-04-07
     16 <150> PRIOR APPLICATION NUMBER: 09/106,377
     17 <151> PRIOR FILING DATE: 1998-06-29
                                                                 Does Not Comply
     19 <150> PRIOR APPLICATION NUMBER: 08/252,508
                                                                 Corrected Diskette Needed
     20 <151> PRIOR FILING DATE: 1994-06-01
                                                                   Cpg-1,2)
    22 <160> NUMBER OF SEQ ID NOS: 71
     24 <170> SOFTWARE: FastSEQ for Windows Version 4.0
ERRORED SEQUENCES
     242 <210> SEQ ID NO: 20
     243 <211> LENGTH: 96
    244 <212> TYPE: DNA
    245 <213> ORGANISM: Artificial Sequence
     247 <220> FEATURE:
     248 <223> OTHER INFORMATION: Primer
    250 <400> SEQUENCE: 20
     251 ctagacgccc tgcagcaaaa gcagggtgac aaagacataa tggagaaaaa aatcactggg 60
   > 252 tataccaccg ttgatatatc ccaatcgcat cgtaaa
    722 <210> SEQ ID NO: 64
723 ______> Pls gnsert C2117 Length: 40
V--> 723 ·___
     724 <212> TYPE: DNA
     725 <213> ORGANISM: Artificial sequence
W--> 727 <220> FEATURE:
    728 <223> OTHER INFORMATION: Oligonucleotide
E--> 730 ·
    730 <400> SEQUENCE: 64
731 tgggtatacc accgttgata tatcccaatc gcatcgtaaa
                                                          40
    733 <210> SEQ ID NO: 65
                                -> found 52
    734 <211> LENGTH: 42
    735 <212> TYPE: DNA
    736 <213> ORGANISM: Artificial sequence
    738 <220> FEATURE:
    739 <223> OTHER INFORMATION: Oligonucleotide
     741 <400> SEQUENCE: 65
E--> 742 gtgatttttt tctccattat gtctttgtca ccctgctttt gctgcagggc gt
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/821,001A

2210> 67
2211> 53
2212> RNA
213> Artificial sequence

3 Noalid Numeric Identifier,
220>
220>
2213> Transcribed M-wt template

400> 67
agcaaaagca gggugaaguu uaaaugauau gaaaaaacac ccuuguuucu acu 53

Scruce of Genetic
WalerialPls See

Bage-3

Jor Error

Expknation

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/821,001A

DATE: 03/03/2006 TIME: 15:07:15

Input Set : A:\NS104D1C1_seqlisting_rev1.txt
Output Set: N:\CRF4\03032006\J821001A.raw

Use of <220> Feature (NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104,pp.29631-32) (Sec.1.823 of new Rules)

Seq#:67

DATE: 03/03/2006

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/821,001A TIME: 15:07:15

Input Set : A:\NS104D1C1_seqlisting_rev1.txt
Output Set: N:\CRF4\03032006\J821001A.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application Number
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:252 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:96 SEQ:20
L:723 M:283 W: Missing Blank Line separator, <220> field identifier
L:723 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:730 M:282 E: Numeric Field Identifier Missing, <211> is required.
L:731 M:252 E: No. of Seq. differs, <211> LENGTH:Input:0 Found:40 SEQ:64
L:742 M:254 E: No. of Bases conflict, LENGTH:Input:42 Counted:52 SEQ:65
L:761 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:763 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:67, <213> ORGANISM:Artificial sequence
L:763 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:67, Line#:763